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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=2; day=19; hr=10; min=14; sec=54; ms=765;]

=====

Reviewer Comments:

<110> Jamal, TEMSAMANI

Michel, KACZOREK

Annik, COLIN DE VERDI?RE

<110> SYNT:EM SA

Please remove the second <110>: it appears only once, next to the first applicant. The question mark in the third applicant's name is due to a foreign accent mark: do not include foreign accent marks: they cannot be processed.

<120> Pharmaceutical composition comprising an anti-cancer agent and at least a peptide

The above <120> response exceeds the Sequence Rules' required 72-character limit.

<140> US 09/856,940

<141> 11-26-1999

<150> FR98/15073

<151> 11-30-1998

<150> PCT/FR99/02939

<151> 11-26-1999

The <141> and <151> responses are in invalid format: use YYYY-MM-DD format.

<160> 13

Although the above <160> response is "13," 17 sequences are in the submitted file.

<210> 1
<211> 18
<212> PRT
<213> porcus

The above <213> response is insufficient: per Sequence Rules, indicate Genus species. This response appears in subsequent sequences.

<210> 8
<211> 16
<212> PRT
<213> unidentified
<220>
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<222> (1)..(16)
<223> X1 to X16 are amino acids of which 6 to 10 of them are hydrophobic amino acids and X6 is Tryptophane
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<223> X4
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<223> X5
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<400>  8

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Xaa Xaa Xaa Xaa Xaa Trp Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

<213> response is invalid: per Sequence Rules, use only: the Genus species, "Artificial Sequence," and "Unknown." "Artificial Sequence" and "Unknown" require explanation in the <220>-<223> section. Give the source of the genetic material. This error appears in subsequent sequences. The explanation for Xaa is very unclear: please state which residues the Xaa's represent. Remove the <220>-<223> sections showing just the location and "X1," for instance.

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<210> 9
<211> 18
<212> PRT
<213> unidentified
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<223> Groups B may be identical or different, and represent a natural
or non-natural amino acid, including amino acid with D configuration and
represent an amino acid for which the side chain carries a basic group
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non-natural amino acid, including amino acid with D configuration and
represent an amino acid for which the side chain carries a basic group
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Besides an invalid <213> response, the explanations for Xaa are again unclear, and the lines too long. "Groups B may be identical or different"--eliminate this--it is very confusing.

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<211> 14
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<210> 17
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<220>
<221> PEPTIDE
<222> (1)..(18)
<400> 17

Arg Gly Gly Arg Leu Ala Tyr Leu Arg Arg Arg Trp Ala Val Leu Val
1 5 10 15

Gly Arg

1

1

The last sequence is shown above. Please remove the "1's" above, which appear at the end of the file.

Application No: 09856940 Version No: 2.0

Input Set:

Output Set:

Started: 2008-02-12 12:58:14.564
Finished: 2008-02-12 12:58:17.414
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 850 ms
Total Warnings: 7
Total Errors: 21
No. of SeqIDs Defined: 13
Actual SeqID Count: 16

Error code	Error Description
E 248	Order Sequence Error <110> -> <110>; Expected Mandatory Tag: <120> in Header
E 287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <141>
E 287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <151>
E 287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <151>
W 402	Undefined organism found in <213> in SEQ ID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (5)
W 402	Undefined organism found in <213> in SEQ ID (6)
W 402	Undefined organism found in <213> in SEQ ID (7)
E 356	Organism is not permitted in <213> in SEQ ID (8)
E 356	Organism is not permitted in <213> in SEQ ID (9)
E 356	Organism is not permitted in <213> in SEQ ID (10)
E 356	Organism is not permitted in <213> in SEQ ID (11)
E 356	Organism is not permitted in <213> in SEQ ID (12)
E 356	Organism is not permitted in <213> in SEQ ID (13)
E 356	Organism is not permitted in <213> in SEQ ID (13)
E 331	Count of Protein differs from the <211> tag Input: 14

Input Set:

Output Set:

Started: 2008-02-12 12:58:14.564
Finished: 2008-02-12 12:58:17.414
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 850 ms
Total Warnings: 7
Total Errors: 21
No. of SeqIDs Defined: 13
Actual SeqID Count: 16

Error code	Error Description
E 212	Invalid Sequence ID Number; Expected 14 as next SeqID but skipped
E 356	Organism is not permitted in <213> in SEQ ID (15)
E 356	Organism is not permitted in <213> in SEQ ID (16)
E 356	Organism is not permitted in <213> in SEQ ID (17)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (17)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (17)
E 252	Calc# of Seq. differs from actual; 13 seqIds defined; count=16

SEQUENCE LISTING

<110> Jamal, TEMSAMANI
 Michel, KACZOREK
 Annik, COLIN DE VERDI?RE
 <110> SYNT:EM SA

<120> Pharmaceutical composition comprising an anti-cancer agent and at least a peptide

<130> 320654US0PCT

<140> US 09/856,940

<141> 11-26-1999

<150> FR98/15073

<151> 11-30-1998

<150> PCT/FR99/02939

<151> 11-26-1999

<160> 13

<170> PatentIn version 3.1

<210> 1

<211> 18

<212> PRT

<213> porcus

<220>

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<400> 1

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Gly Arg

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<213> porcus

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Cys Arg

<210> 8
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 Tryptophane
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chain carries a basic group
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<223> Groups X may be identical or different, and represent a natural or non-natural amino acid, including amino acid with D configuration and represent a residue of aliphatic or aromatic amino acid
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Xaa Asx

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<220>

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amino acid
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Asx

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Arg

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<213> unidentified
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<213> unidentified
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1 5 10 15

Gly Arg

1

1